

K. LACOURCIERE

1635

#12

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/08/765,244

DATE: 05/04/2000  
TIME: 12:18:32

Input Set : A:\8484-018.txt  
Output Set: C:\CRF3\REFHOLD\05042000\H765244.raw

4 <110> APPLICANT: Seibel, Peter  
5 Seibel, Andrea  
7 <120> TITLE OF INVENTION: CHIMERICAL PEPTIDE-NUCLEIC ACID  
8 FRAGMENT, PROCESS FOR PRODUCING THE SAME AND ITS USES FOR  
9 APPROPRIATELY INTRODUCING NUCLEIC ACIDS INTO CELL ORGANELLES  
10 AND CELLS  
12 <130> FILE REFERENCE: 8484-0018-999  
14 <140> CURRENT APPLICATION NUMBER: US 08/765,244  
15 <141> CURRENT FILING DATE: 1997-10-30  
17 <150> PRIOR APPLICATION NUMBER: PCT/DE95/00775  
18 <151> PRIOR FILING DATE: 1995-06-11  
20 <150> PRIOR APPLICATION NUMBER: DE P 44 21 079.5  
21 <151> PRIOR FILING DATE: 1994-06-16  
23 <160> NUMBER OF SEQ ID NOS: 22  
25 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
27 <210> SEQ ID NO: 1  
28 <211> LENGTH: 41  
29 <212> TYPE: PRT  
30 <213> ORGANISM: Rattus rattus  
32 <220> FEATURE:  
34 <400> SEQUENCE: 1  
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36 1 5 10 15  
37 Ala His Thr Ser Met Val Arg Asn Phe Arg Tyr Gly Lys Pro Val Gln  
38 20 25 30  
39 Ser Gln Leu Lys Pro Arg Asp Leu Cys  
40 35 40  
43 <210> SEQ ID NO: 2  
44 <211> LENGTH: 39  
45 <212> TYPE: DNA  
46 <213> ORGANISM: Artificial Sequence  
48 <220> FEATURE:  
49 <223> OTHER INFORMATION: Synthetic sequence  
51 <400> SEQUENCE: 2  
52 ccccggttac cctgcgagcc ctgggctcgc aaggtaccc  
54 <210> SEQ ID NO: 3  
55 <211> LENGTH: 34  
56 <212> TYPE: DNA  
57 <213> ORGANISM: Homo sapiens  
59 <400> SEQUENCE: 3  
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62 <210> SEQ ID NO: 4  
63 <211> LENGTH: 34  
64 <212> TYPE: DNA  
65 <213> ORGANISM: Homo sapiens  
67 <400> SEQUENCE: 4  
68 ccatactact aatctcatca atagagaggg cccc

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72 <212> TYPE: DNA
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76 ccatactact aatctcatca atagagaggg 30
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79 <211> LENGTH: 103
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Chimeric sequence
86 <400> SEQUENCE: 6
87 ccatactact aatctcatca atagagaggg ccccggttac ctgcgagcc ctgggctcgc 60
88 aagggtaccg gggccctctc tattgatgag attagtagta tgg 103
90 <210> SEQ ID NO: 7
91 <211> LENGTH: 1465
92 <212> TYPE: DNA
93 <213> ORGANISM: Homo sapiens
95 <400> SEQUENCE: 7
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97 gtttattgct aaaggttaat cactgctgtt tcccggtggg gtgtggctag gctaagcgtt 120
98 ttgagctgca ttgctgcgtg cttgatgctt gtcccttttg atcgtgggtga tttagagggt 180
99 gaactcactg gaacggggat gcttgcattgt gtaatcttac taagagctaa tagaaaggct 240
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103 agattagtag tatgggagtg ggaggggaaa ataattgtgt agttgggggg tgactgttaa 480
104 aagtgcatac cgccaaaaga taaaatttga aatctgttta ggctgggtgt agggcccttt 540
105 gtcccacacc caccacaaga cagggtttgt taagatggca gagcccgta atgcataaa 600
106 acttaaaact ttacagtcag aggttcaatt cctcttctta acaacatacc catggccaac 660
107 ctctactctc tcattgtacc cattctaatc gcaatggcat tctaatgct taccgaacga 720
108 aaaattctag gctatataca actacgcaaa ggccccaacg tggtagggcc ctacgggcta 780
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114 ctagccatca ttctactatc aacattacta ataagtggtc cctttaacct ctccaccctt 1140
115 atcacaacac aagaacacct ctgattactc ctgccatcat gacccttggc cataaatatg 1200
116 ttatctcca cactagcaga gaccaaccga acccccttcg accttgccga aggggagtc 1260
117 gaactagctc caggcttcaa catcgaatac gcgcaggcc ccttcgccct attcttcata 1320
118 gccgaataca caaacattat tataataaac accctacca ctacaattct cctaggaaca 1380
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120 ctaacctccc tgttcttatg aattc 1465
122 <210> SEQ ID NO: 8
123 <211> LENGTH: 1465
124 <212> TYPE: DNA
125 <213> ORGANISM: Homo sapiens

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130 ttataataat gtttgtgtat tcggtctatga agaatagggc gaaggggcct gcggcgatt 180
131 cgatgttgaa gcctgagact agttcggact ccccttcggc aaggtcgaag ggggttcggt 240
132 tggctctctgc tagtgtggag ataaatcata ttatggccaa gggctcatgat ggcaggagta 300
133 atcagaggtg ttcttgtgtt gtgataaggg tggagaggtt aaaggagcca cttattagta 360
134 atgttgatag tagaatgatg gctagggtga ctccatata gattgttttg gctactgctc 420
135 gcagtgcgcc gatcagggcg tagtttgagt ttgatgctca cctgatcag aggattgagt 480
136 aaacggctag gctagaggtg gctagaataa ataggagccc taggttgagg ttgaccaggg 540
137 ggttggttat ggggaggggg gttcatagta gaagagcgat ggtgagagct aaggtcgggg 600
138 cgtgatgta gaggggtgat gtagatgtgg cgggttttag gggctctttg gtgaagagt 660
139 ttatggcgctc agcgaagggg ttagtagacc cgtaggggcc taccacgttg gggcctttgc 720
140 gtatgtgtat atagcctaga atttttcggt cggttaagcat taggaatgcc attgcgatta 780
141 gaatgggtac aatgaggagt aggaggttgg ccatgggtat gttgttaaga agagggaattg 840
142 aacctctgac tgtaaagttt taagttttat gcgattaccg ggctctgcca tcttaacaaa 900
143 cctgtttctt ggggtgggtg gggacaaagg gccctaacac cagcctaacc agatttcaaa 960
144 ttttatcttt tggcggtatg cacttttaac agtcaccccc caactaacac attattttoc 1020
145 cctcccactc ccatactact aatctcatca atacaacccc cgcctatcct acccagcaca 1080
146 cacacaccgc tgctaacccc ataccgccga ccaaccaaac cccaagaca cccccacag 1140
147 ttatgttagc ttacctctc aaagcaatac actgaaaatg tttagacggg ctcacatcac 1200
148 ccataaaca aataggtttg gtctagcct ttctattagc tcttagtaag attacacatg 1260
149 caagcatccc cggtccagtg agttcaccct ctaaatcacc acgatcaaaa ggaacaagca 1320
150 tcaagcacgc agcaatgcag ctcaaaacgc tttagcctagc cacaccccca cgggaaacag 1380
151 cagtgattaa cctttagcaa taaacgaaag tttaactaag ctatactaac cccaggggtg 1440
152 gtcaatttcg tgccagccac cgcgg 1465
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155 <211> LENGTH: 86
156 <212> TYPE: DNA
157 <213> ORGANISM: Artificial Sequence
159 <220> FEATURE:
160 <223> OTHER INFORMATION: Synthetic sequence
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164 cccggtaac gcataaaatg agaccg 86
166 <210> SEQ ID NO: 10
167 <211> LENGTH: 94
168 <212> TYPE: DNA
169 <213> ORGANISM: Artificial Sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: Synthetic sequence
174 <400> SEQUENCE: 10
175 gatccggtct cattttatgc gattaccggg ctctgccatc ttaacaaatg atcatatggt 60
176 gacctgaggc gccttaagct tcgcgatatc tgca 94
178 <210> SEQ ID NO: 11
179 <211> LENGTH: 1455
180 <212> TYPE: DNA
181 <213> ORGANISM: Unknown
183 <220> FEATURE:

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Input Set : A:\8484-018.txt  
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188 tgtgagcccg tctaaacatt ttcagtgtat tgctttgagg aggtaagcta cataaactgt 120
189 gggggggtgc tttgggggtt ggttggttcg gggatatggg ttagcagcgg tgtgtgtgtg 180
190 ctgggttaga tggcgggggg ttgtattgat gagattagta gtatgggagt gggaggggaa 240
191 aataatgtgt tagttggggg gtgactgtta aaagtgcata ccgccaaaag ataaaatttg 300
192 aaatctgtgt aggtcgtgtg taggggttctt tgtttttggg gtttggcaga gatgtgttta 360
193 agtgctgtgg ccagaagcgg gggagggggg gtttgggtga aattttttgt tatgatgtct 420
194 gtgtggaag tggtgtgtca gacattcaat tgttattatt atgtcctaca agcattaatt 480
195 aattaacaca ctttagtaag tatgttcgcc tgaatattg aacgtagggt cgataaataa 540
196 taggatgagg caggaatcaa agacagatac tgcgacatag ggtgctccgg ctccagcgtc 600
197 tcgcaatgct atcgcgtgtca tccccccag acgaaaatac caaatgcatg gagagctccc 660
198 gtgagtgggt aatagggtga tagacctgtg atccatcgtg atgtcttatt taaggggaac 720
199 gtgtgggcta tttaggcttt atgacctga agtaggaacc agatgtcggg tacagttcac 780
200 tttagctacc cccaagtgtt atgggcccgg agcgaggaga gtgacactct tgtgcgggat 840
201 attgatttca cggaggatgg tggtaaggg acccctatct gaggggggtc atccatgggg 900
202 acgagaaggg atttactgt aatgtgctat gtacggtaaa tggctttatg tactatgtac 960
203 tgttaagggt ggttaggttt gttggtatcc tagtgggtga ggggtggcct tggagttgca 1020
204 gttgatgtgt gatagttgag ggttgattgc tgtacttgct tgaagcatg gggagggggt 1080
205 tttgatgtgg attgggtttt tatgtactac aggtgggtcaa gtatttatgg taccgtacaa 1140
206 tattcatggt ggtcggcagt aatgtacgaa atacatagcg gttgtgatg ggtgagtcga 1200
207 tacttgggtg gtaccctaat ctgcttcccc atgaagaac agagaatag ttaaattaga 1260
208 atcttagctt tgggtgtcaa tgggtgagtt aaagactttt tctctgattt gtcottggaa 1320
209 aaagggtttc atctcgggtt tacaagactg gtgtattagc tgcagatata gcgaagctta 1380
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211 aatgagaccg gatcc 1455
213 <210> SEQ ID NO: 12
214 <211> LENGTH: 1455
215 <212> TYPE: DNA
216 <213> ORGANISM: Unknown
218 <220> FEATURE:
219 <223> OTHER INFORMATION: Plasmid
221 <400> SEQUENCE: 12
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224 gagatgaaaa cttttttcca aggacaaatc agagaaaaag tctttaactc caccattagc 180
225 acccaaaact aagattctaa tttaaactat tctctgttct ttcattggga agcagatttg 240
226 ggtaccacc cagatttgac tcacccatca acaaccgcta tgtatttcgt acattactgc 300
227 cagccaccat gaattattga cggtagcata aatacttgac cacctgtagt acataaaaac 360
228 ccaatccaca tcaaaacccc ctcccatgc ttacaagcaa gtacagcaat caacctcaa 420
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232 ctccgtgaaa tcaatatccc gcacaagagt gctactctcc tcgctcgggg cccataaac 660
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234 ctaaaatagc cacacgttcc ccttaaataa gacatcacga tggatcacag gtcattacac 780
235 ctattaacca ctacgggag ctctccatgc atttggtatt ttcgtctggg gggtagtcac 840
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237 tctgcctca tcctattatt tatgcacct acgttcaata ttacaggcga acatacttac 960
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239 agccacttcc cacacagaca tcataacaaa aaatttccac caaaccccc ctccccgcgt 1080
240 tctggccaca gcacttaaac acatctctgc caaaccccaa aaacaaagaa ccctaacacc 1140
241 agcctaacca gatttcaaat tttatctttt ggcgggtatgc acttttaaca gtcaccccc 1200
242 aactaacaca ttattttccc ctccactcc cactactacta atctcatcaa tacaaccccc 1260
243 gccatccta cccagcacac acacacgcgt gctaacccca taccgccgaac caaccaaacc 1320
244 ccaaagacac cccccacagt ttatgtagct tacctctca aagcaataca ctgaaaatgt 1380
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259 ggggggtgtc tttgggtttt ggttggttcg gggtagggg ttagcagcgg tgtgtgtgtg 180
260 ctgggttaga tgggcggggg ttgtattgat gagattagta gtatgggagt gggaggggaa 240
261 aataatgtgt tagttggggg gtgactgtta aaagtgcata ccgccaaaag ataaaatttg 300
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263 agtctgttgg ccagaagcgg gggagggggg gtttggttga aattttttgt tatgatgtct 420
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270 tttagctacc cccaagtgtt atgggcccgg agcaggagga gtagcactct tgtgcgggat 840
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273 tgttaagggt gggtaggttt gttgggtacc tagtgggtga ggggtggcgt tggagttgca 1020
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287 tagaggcgac aaacctacgg agcctggtga tagctggttg tccaagatag aatcttagtt 1860
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